

## **RAW SEQUENCE LISTING**

**The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.**

Application Serial Number: US/10/123,383

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RAW SEQUENCE LISTING  
PATENT APPLICATION: US/10/723,383

DATE: 12/20/2004  
TIME: 12:05:08

Input Set : A:\-29-2.app  
Output Set: N:\CRF4\12202004\J723383.raw

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3 <110> APPLICANT: Nasoff, Marc
4     Deveraux, Quinn L.
5     Knee, Deborah A.
6     Aza-Blanc, Pedro
7     Hampton, Garret M.
8     Wagner, Klaus
9     IRM LLC
11 <120> TITLE OF INVENTION: Methods and Compositions for Inducing Apoptosis in
12     Cancer Cells
14 <130> FILE REFERENCE: 021288-002920US
16 <140> CURRENT APPLICATION NUMBER: US 10/723,383
17 <141> CURRENT FILING DATE: 2003-11-25
19 <150> PRIOR APPLICATION NUMBER: US 60/429,842
20 <151> PRIOR FILING DATE: 2002-11-27
22 <150> PRIOR APPLICATION NUMBER: US 448,960
23 <151> PRIOR FILING DATE: 2003-02-21
25 <150> PRIOR APPLICATION NUMBER: US 60/494,714
26 <151> PRIOR FILING DATE: 2003-08-12
28 <150> PRIOR APPLICATION NUMBER: US 60/504,901
29 <151> PRIOR FILING DATE: 2003-09-22
31 <160> NUMBER OF SEQ ID NOS: 15
33 <170> SOFTWARE: PatentIn Ver. 2.1
35 <210> SEQ ID NO: 1
36 <211> LENGTH: 64
37 <212> TYPE: PRT
38 <213> ORGANISM: Artificial Sequence
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41 <223> OTHER INFORMATION: Description of Artificial Sequence:Baculovirus
42     Inhibitory Repeat (BIR) region motif conserved
43     residue consensus sequence
45 <220> FEATURE:
46 <221> NAME/KEY: MOD_RES
47 <222> LOCATION: (2)..(24)
48 <223> OTHER INFORMATION: Xaa = any amino acid, Xaa at positions 22-24 may
49     be present or absent
51 <220> FEATURE:
52 <221> NAME/KEY: MOD_RES
53 <222> LOCATION: (26)..(36)
54 <223> OTHER INFORMATION: Xaa = any amino acid
56 <220> FEATURE:
57 <221> NAME/KEY: MOD_RES
58 <222> LOCATION: (38)..(39)
59 <223> OTHER INFORMATION: Xaa = any amino acid

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61 <220> FEATURE:  
 62 <221> NAME/KEY: MOD\_RES  
 63 <222> LOCATION: (41)..(56)  
 64 <223> OTHER INFORMATION: Xaa = any amino acid  
 66 <220> FEATURE:  
 67 <221> NAME/KEY: MOD\_RES  
 68 <222> LOCATION: (58)..(63)  
 69 <223> OTHER INFORMATION: Xaa = any amino acid  
 71 <400> SEQUENCE: 1  
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 73 1 5 10 15  
 75 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Gly Xaa Xaa Xaa Xaa Xaa Xaa  
 76 20 25 30  
 78 Xaa Xaa Xaa Xaa Cys Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa  
 79 35 40 45  
 81 Xaa Xaa Xaa Xaa Xaa Xaa Xaa His Xaa Xaa Xaa Xaa Xaa Cys  
 82 50 55 60  
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 86 <211> LENGTH: 359  
 87 <212> TYPE: DNA  
 88 <213> ORGANISM: Artificial Sequence  
 90 <220> FEATURE:  
 91 <223> OTHER INFORMATION: Description of Artificial Sequence:anti-DR5  
 92 Antibody A light chain variable region  
 94 <400> SEQUENCE: 2  
 95 gacattgcga tgacccagtc tcacaagttc atgtccacat tagtggaga cagggtcagc 60  
 96 atcacctgca aggccagtca ggatgtaat actgctatag cctggtatca acaaaaacca 120  
 97 gggcaatctc ctaaactact gatttactgg gcatccaccc ggcacactgg agtccctgtat 180  
 98 cgcttcacag gcagtggatc tgggacagat tatactctca ccatcagcag tatggaggct 240  
 99 gaagatgctg ccacttatta ctgccagcag tggagtagta acccgctcac gttcggtgct 300  
 100 gggaccaagc tggagctgaa acgggctgat gctgcaccaa ctgtatccat cttcccacc 359  
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 104 <211> LENGTH: 360  
 105 <212> TYPE: DNA  
 106 <213> ORGANISM: Artificial Sequence  
 108 <220> FEATURE:  
 109 <223> OTHER INFORMATION: Description of Artificial Sequence:anti-DR5  
 110 Antibody A heavy chain variable region  
 112 <400> SEQUENCE: 3  
 113 caggcaaagg tccagctgca gcagtctgga gctgagctgg tggaaacccgg ggcacatcagt 60  
 114 aagctgtcct gcaaggcttc tggotacacc ttcaactgact atactataca ctgggtaaag 120  
 115 cagaggtctg gacagggtct tgagtggatt gggtggttt accctggagg tggttatata 180  
 116 aaatacaatg agaaaattcaa ggacaggccc acattgactg cggacaaaatc ctccaacaca 240  
 117 gtctatatgg agcttagtgc attgacatct gaaggctctg cggcttattt ctgtgcaaga 300  
 118 cacgaagagg gcatcttattt tgactactgg ggccaaggca ccactctcac agtctcctca 360  
 121 <210> SEQ ID NO: 4  
 122 <211> LENGTH: 118  
 123 <212> TYPE: PRT  
 124 <213> ORGANISM: Artificial Sequence

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126 <220> FEATURE:  
 127 <223> OTHER INFORMATION: Description of Artificial Sequence:anti-DR5  
 128       Antibody A heavy chain subgroup B variable region  
 130 <400> SEQUENCE: 4  
 131 Lys Val Gln Leu Gln Gln Ser Gly Ala Glu Leu Val Lys Pro Gly Ala  
 132       1               5               10               15  
 134 Ser Val Lys Leu Ser Cys Lys Ala Ser Gly Tyr Thr Phe Thr Asp Tyr  
 135       20              25              30  
 137 Thr Ile His Trp Val Lys Gln Arg Ser Gly Gln Gly Leu Glu Trp Ile  
 138       35              40              45  
 140 Gly Trp Phe Tyr Pro Gly Gly Tyr Ile Lys Tyr Asn Glu Lys Phe  
 141       50              55              60  
 143 Lys Asp Arg Ala Thr Leu Thr Ala Asp Lys Ser Ser Asn Thr Val Tyr  
 144       65              70              75              80  
 146 Met Glu Leu Ser Arg Leu Thr Ser Glu Gly Ser Ala Val Tyr Phe Cys  
 147       85              90              95  
 149 Ala Arg His Glu Glu Gly Ile Tyr Phe Asp Tyr Trp Gly Gln Gly Thr  
 150       100             105             110  
 152 Thr Leu Thr Val Ser Ser  
 153       115  
 156 <210> SEQ ID NO: 5  
 157 <211> LENGTH: 109  
 158 <212> TYPE: PRT  
 159 <213> ORGANISM: Artificial Sequence  
 161 <220> FEATURE:  
 162 <223> OTHER INFORMATION: Description of Artificial Sequence:anti-DR5  
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 164       region  
 166 <400> SEQUENCE: 5  
 167 Asp Ile Ala Met Thr Gln Ser His Lys Phe Met Ser Thr Leu Val Gly  
 168       1              5              10              15  
 170 Asp Arg Val Ser Ile Thr Cys Lys Ala Ser Gln Asp Val Asn Thr Ala  
 171       20              25              30  
 173 Ile Ala Trp Tyr Gln Gln Lys Pro Gly Gln Ser Pro Lys Leu Leu Ile  
 174       35              40              45  
 176 Tyr Trp Ala Ser Thr Arg His Thr Gly Val Pro Asp Arg Phe Thr Gly  
 177       50              55              60  
 179 Ser Gly Ser Gly Thr Asp Tyr Thr Leu Thr Ile Ser Ser Met Glu Ala  
 180       65              70              75              80  
 182 Glu Asp Ala Ala Thr Tyr Tyr Cys Gln Gln Trp Ser Ser Asn Pro Leu  
 183       85              90              95  
 185 Thr Phe Gly Ala Gly Thr Lys Leu Glu Leu Lys Arg Ala  
 186       100             105  
 189 <210> SEQ ID NO: 6  
 190 <211> LENGTH: 23  
 191 <212> TYPE: DNA  
 192 <213> ORGANISM: Artificial Sequence  
 194 <220> FEATURE:  
 195 <223> OTHER INFORMATION: Description of Artificial Sequence:synthetic siRNA

**RAW SEQUENCE LISTING  
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196 ttds (N) 19TT  
 198 <220> FEATURE:  
 199 <221> NAME/KEY: modified\_base  
 200 <222> LOCATION: (3)..(21)  
 201 <223> OTHER INFORMATION: n = any nucleotide  
 203 <400> SEQUENCE: 6  
**204 ttnnnnnnnn nnnnnnnnnn ntt** 23  
 207 <210> SEQ ID NO: 7  
 208 <211> LENGTH: 354  
 209 <212> TYPE: DNA  
 210 <213> ORGANISM: Artificial Sequence  
 212 <220> FEATURE:  
 213 <223> OTHER INFORMATION: Description of Artificial Sequence:alternate  
 214 sequence for anti-DR5 Antibody A heavy chain  
 215 variable region  
 217 <400> SEQUENCE: 7  
 218 aaggtccagc tgcagcagtc tggagctgag ctggtaaaac cggggcattt aatgtttttttt 60  
 219 tcctgcaagg cttctggcta caccttcaact gactatacta tacactgggt aaagcagagg 120  
 220 tctggacagg gtcttgatgt gattgggtgg ttttacccctg gaggtggta tataaaaatac 180  
 221 aatgagaaat tcaaggacag gcccacattt actgcggaca aatcctccaa cacagtctat 240  
 222 atggagctt aatggatggac atctgaagac tctgcggctt atttctgtgc aagacacgaa 300  
 223 gagggcatct attttactt ctggggccaa ggcaccactc tcacagtctc ctca 354  
 226 <210> SEQ ID NO: 8  
 227 <211> LENGTH: 118  
 228 <212> TYPE: PRT  
 229 <213> ORGANISM: Artificial Sequence  
 231 <220> FEATURE:  
 232 <223> OTHER INFORMATION: Description of Artificial Sequence:alternate  
 233 sequence for anti-DR5 Antibody A heavy chain  
 234 variable region  
 236 <400> SEQUENCE: 8  
 237 Lys Val Gln Leu Gln Gln Ser Gly Ala Glu Leu Val Lys Pro Gly Ala  
 238 1 5 10 15  
 240 Ser Val Lys Leu Ser Cys Lys Ala Ser Gly Tyr Thr Phe Thr Asp Tyr  
 241 20 25 30  
 243 Thr Ile His Trp Val Lys Gln Arg Ser Gly Gln Gly Leu Glu Trp Ile  
 244 35 40 45  
 246 Gly Trp Phe Tyr Pro Gly Gly Tyr Ile Lys Tyr Asn Glu Lys Phe  
 247 50 55 60  
 249 Lys Asp Arg Ala Thr Leu Thr Ala Asp Lys Ser Ser Asn Thr Val Tyr  
 250 65 70 75 80  
 252 Met Glu Leu Ser Arg Leu Thr Ser Glu Asp Ser Ala Val Tyr Phe Cys  
 253 85 90 95  
 255 Ala Arg His Glu Glu Gly Ile Tyr Phe Asp Tyr Trp Gly Gln Gly Thr  
 256 100 105 110  
 258 Thr Leu Thr Val Ser Ser  
 259 115  
 262 <210> SEQ ID NO: 9  
 263 <211> LENGTH: 312

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264 <212> TYPE: DNA  
 265 <213> ORGANISM: Artificial Sequence  
 267 <220> FEATURE:  
 268 <223> OTHER INFORMATION: Description of Artificial Sequence:alternate  
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 270 variable region  
 272 <400> SEQUENCE: 9  
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 274 atcacctgca aggccagtc ggtatgtaat actgtctatag cctgttatca acaaaaacca 120  
 275 gggcaatctc ctaaactact gatttactgg gcattccaccc ggcacactgg agtccctgtat 180  
 276 cgcttcacag gcagtggatc tgggacagat tatactctca ccatcagcag tgtgcaggct 240  
 277 gaagacctgg cactttatta ctgtcagcaa cattatacca ctccattcac gttcggctcg 300  
 278 gggacaaaatg 312  
 281 <210> SEQ ID NO: 10  
 282 <211> LENGTH: 104  
 283 <212> TYPE: PRT  
 284 <213> ORGANISM: Artificial Sequence  
 286 <220> FEATURE:  
 287 <223> OTHER INFORMATION: Description of Artificial Sequence:alternate  
 288 sequence for anti-DR5 Antibody A light chain  
 289 variable region  
 291 <400> SEQUENCE: 10  
 292 Asp Ile Val Met Thr Gln Ser His Lys Phe Met Ser Thr Ser Val Gly  
 293 1 5 10 15  
 295 Asp Arg Val Ser Ile Thr Cys Lys Ala Ser Gln Asp Val Asn Thr Ala  
 296 20 25 30  
 298 Ile Ala Trp Tyr Gln Gln Lys Pro Gly Gln Ser Pro Lys Leu Leu Ile  
 299 35 40 45  
 301 Tyr Trp Ala Ser Thr Arg His Thr Gly Val Pro Asp Arg Phe Thr Gly  
 302 50 55 60  
 304 Ser Gly Ser Gly Thr Asp Tyr Thr Leu Thr Ile Ser Ser Val Gln Ala  
 305 65 70 75 80  
 307 Glu Asp Leu Ala Leu Tyr Tyr Cys Gln Gln His Tyr Thr Thr Pro Phe  
 308 85 90 95  
 310 Thr Phe Gly Ser Gly Thr Lys Leu  
 311 100  
 314 <210> SEQ ID NO: 11  
 315 <211> LENGTH: 19  
 316 <212> TYPE: DNA  
 317 <213> ORGANISM: Artificial Sequence  
 319 <220> FEATURE:  
 320 <223> OTHER INFORMATION: Description of Artificial Sequence:siPAK1-0 siRNA  
 321 directed against PAK1  
 323 <400> SEQUENCE: 11  
 324 agagctgcta cagcatcaa 19  
 327 <210> SEQ ID NO: 12  
 328 <211> LENGTH: 19  
 329 <212> TYPE: RNA  
 330 <213> ORGANISM: Artificial Sequence

RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/10/723,383

DATE: 12/20/2004  
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Input Set : A:\-29-2.app  
Output Set: N:\CRF4\12202004\J723383.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; Xaa Pos. 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23  
Seq#:1; Xaa Pos. 24, 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 39, 40, 41, 42, 43, 44, 45  
Seq#:1; Xaa Pos. 46, 47, 48, 49, 50, 51, 52, 53, 54, 55, 56, 57, 58, 59, 60, 61, 62, 63  
Seq#:6; N Pos. 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21

**VERIFICATION SUMMARY** DATE: 12/20/2004  
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Input Set : A:\-29-2.app  
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L:72 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0

M:341 Repeated in SeqNo=1

L:204 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:0